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- Applicant: SCHERING CORPORATION
 2000 Gelloping Hill Road
 Kenilworth New Jersey 07033(US)
- (2) Inventor: Galizzi, Jean-Pierre 27 chemin des peupliers BP11 F-69572 Dardilly(FR) Inventor: Harada, Nobuyuki 2310 St. Francis Drive Palo Alto, California 94303(US) Inventor: Miyajima, Atsushi 4159 Drake Avenue Palo Alto, California 94306(US)
- Representative: Ritter, Stephen David et al. Mathys & Squire 10 Fleet Street London EC4Y 1AY(GB)

(9) Interleukin-4 receptors.

Nucleic acids are provided which encode mammalian interleukin-4 receptors, and which serve as probes for related receptors by cross hybridization. Soluble forms of human interleukin-4 receptors are useful as antagonists of interleukin-4 in the treatment of conditions associated with excessive IgE production.

MAMMALIAN INTERLEUKIN-4 RECEPTOR + USED AS

ANTAGONISTS OF INTERLEUKIN-4 IN TRATING CONDITIONS

ASSOCIATED WITH EXCESS IGE PRODUCTION INCLUDING

ALLERGIC CONDITIONS

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INTERLEUKIN-4 RECEPTORS

Field of the Invention

The invention relates generally to the mammalian IL-4 (interleukin-4) receptor, and more particularly to the synthesis of the IL-4 receptor by recombinant cells and to the manufacture and use of such receptors for screening agonists and antagonists and for reducing in vivo levels of IL-4 in certain disease conditions.

BACKGROUND

Several forms of mammalian IL-4 have been cloned and characterized: e.g. Lee et al.; Proc. Natl. Acad. Sci. USA., Vol. 83, pgs. 2061-2065 (1986); and Yokota et al., Proc. Natl. Acad. Sci. USA., Vol. 83, pgs. 5894-5898 (1988). More recently, studies have shown that IL-4 may play an important role in hypersensitivity reactions, particularly in those due to immunoglobulin E (IgE): e.g. Coffman et al., J. Immunol., Vol. 4538-4541 (1986); Finkelman et al., Proc. Natl. Acad. Sci. USA., Vol. 83, pgs. 9675-9678 (1988); Snapper et al., Science, Vol. 236, pgs. 944-947 (1987); and Mosmann et al., Ann. Rev. Immunol., Vol. 7, pgs. 145-173 (1989). In particular, Finkelman et al. (cited above) have shown that IgE responses can be suppressed by the administration of antibodies capable of blocking the biological activity of IL-4, thereby suggesting an alternative route for treating allergic conditions characterized by excessive IgE production.

Currently, glucocorticoid steroids are the most effective drugs for treating such disorders. However, prolonged steroid treatment is associated with many deleterious side effects (Goodman and Gillman, The Pharmacological Basis of Therapeutics 6th Ed. (MacMillan, New York, 1980)). Accordingly, the availability of alternative approaches to the treatment of immune disorders associated with excessive IgE production could have important clinical utility.

SUMMARY OF THE INVENTION

The invention is directed to mammalian IL-4 receptors ("IL-4Rs") in soluble and in membrane-bound forms, nucleic acids encoding the same, and the use of soluble forms of the receptors ("sIL-4Rs") as antagonists of IL-4 to treat conditions characterized by excessive IgE production. The invention also includes the nucleic acids (1) that are effectively homologous to the murine nucleotide sequence defined by Formula I below, and (2) that encode a polypeptide capable of binding IL-4 derived from the same species as the nucleic acid. Preferably, the soluble IL-4Rs bind IL-4 with substantially the same affinity as the native membrane-bound IL-4Rs, or with even greater affinity.

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	Q1105	55		
	5'-	GCGCGGCGTG	GAGCCTGAAC	TCGCAGGTTC
	TGGCTGGACT	TCTCGAAGCT	GAGGAGAAGC	AGAGGGACCT
	GGCTTCTGAT	TTTGGATCTG	CGTGCTTGCT	GGTTCTGGCG
5	CCTGCTGGTC	TTCTTCCTGT	AACCTAGGAC	TCGGGGCTTG
	CACATGCTTT	TTTTTTGAAG	TTGCTGGAGA	GGGAGCCCAG
	GACCTTGTGC	AGGCACCTTT	TGTGTCCCCA	ATGGGGCGGC
		GTTCCTGACC	TCTGTGGGCT	GTCTGATTTT
10	TTTGCACCAA	ACTGGATCTG	GGAGCATCAA	SGTCCTGGGT
	GCTGTTGGTG	GCTTCTCTGA	CTACATCCGC	ACTTCCACGT
	GAGCCCACCT		GCTGTGGACT	GCAGTTCTCA
15	GTGAGTGGTT	CCTGGATAGC		CGAGTTCTCT
	GCTCTGCCTA	CACTACAGGC	TGATGTTCTT	
	GAAAACCTCA	CATGCATCCC	GAGGAACAGT	GCCAGCACTG
	TGTGTGTGTG	CCACATGGAA	ATGAATAGGC	CGGTCCAATC
20	AGACAGATAC	CAGATGGAAC	TGTGGGCTGA	GCACAGACAG
	CTGTGGCAGG	GCTCCTTCAG	CCCCAGTGGT	AATGTGAAGC
	CCCTAGCTGG	AGACAACCTC	ACACTCCACA	CCAATGTGTC
	CGACGAATGG	CTGCTGACCT	GGAATAACCT	GTACCCATUG
25	CGACGAAIGG	01001011001	-	

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	AACAACTTAC	TGTACAAAGA	CCTCATCTCC	ATGGTCAACA
	TCTCCAGAGA	GGACAACCCT	GCAGAATTCA	TAGTCTATAA
	TGTGACCTAC	AAGGAACCCA	GGCTGAGCTT	CCCGATCAAC
5	ATCCTGATGT	CAGGGGTCTA	CTATACGGCG	CGTGTGAGGG
	TCAGATCCCA	GATACTCACT	GGCACCTGGA	GTGAGTGGAG
	TCCTAGCATC	ACGTGGTACA	ACCACTTCCA	GCTGCCCCTG
10	ATACAGCGCC	TTCCACTGGG	GGTCACCATC	TCCTGCCTCT
	GCATCCCGTT	GTTTTGCCTG	TTCTGTTACT	TCAGCATTAC
	CAAGATTAAG	AAGATATGGT	GGGACCAGAT	TCCCACCCCA
15	GCACGCAGTC	CCTTGGTGGC	CATCATCATT	CAGGATGCAC
,,	AGGTGCCCCT	CTGGGATAAG	CAGACCCGAA	GCCAGGAGTC
	AACCAAGTAC	CCGCACTGGA	AAACTTGTCT	AGACAAGCTG
	CTGCCTTGCT	TGCTGAAGCA	CAGAGTAAAG	AAGAAGACAG
20	ACTTCCCGAA	GGCTGCCCCA	ACCAAGTCTC	TCCAGAGTCC
	TGGAAAGGCA	GGCTGGTGTC	CCATGGAGGT	CAGCAGGACC
	GTCCTCTGGC	CAGAGAATGT	TAGTGTCAGT	GTGGTGCGCT
25	GTATGGAGCT	GTTTGAGGCC	CCAGTACAGA	ATGTGGAGGA
	GGAAGAAGAT	GAGATAGTCA	AAGAGGACCT	GAGCATGTCA
	CCTGAGAACA	GCGGAGGCTG	CGGCTTCCAG	GAGAGCCAGG
30	CAGACATCAT	GGCTCGGCTC	ACTGAGAACC	TGTTTTCCGA
	CTTGTTGGAG	GCTGAGAATG	GGGGCCTTGG	CCAGTCAGCC
	TTGGCAGAGT	CATGCTCCCC	TCTGCCTTCA	GGAAGTGGGC
	AGGCTTCTGT	ATCCTGGGCC	TGCCTCCCCA	TGGGGCCCAG
36	TGAGGAGGCC	ACATGCCAGG	TCACAGAGCA	GCCTTCACAC
-	CCAGGCCCTC	TTTCAGGCAG	CCCAGCCCAG	AGTGCACCTA
	CTCTGGCTTG	CACGCAGGTC	CCACTTGTCC	TTGCAGACAA
40	TCCTGCCTAC	CGGAGTTTTA	GTGACTGCTG	TAGCCCGGCC
	CCAAATCCTG	GAGAGCTGGC	TCCAGAGCAG	CAGCAGGCTG
	ATCATCTGGA	AGAAGAGGAG	CCTCCAAGCC	CGGCTGACTC
	CCATTCTTCA	GGGCCTCCAA	TGCAGCCAGT	GGAGAGCTGG
.45	GAGCAGATCC	TTCACATGAG	TGTCCTGCAG	CATGGGGCAG
	CTGCTGGCC	-31		

Formula 1

In Formula I, A, C, G, and T represent deoxyadenosine, deoxycytidine, deoxyguanosine, and thymidine, respectively. Standard notation is used below to designate L-amino acids: e.g. Cohn, Meth. Enzymol., Vol. 108, pg. 4 (1984).

Brief Description of the Drawings

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Figure 1 illustrates data describing the ability of mouse sIL-4R to block the binding of IL-4 to HT-2 cells. Figure 2 illustrates data describing the ability of mouse sIL-4R to inhibit the biological activity of IL-4.

Figure 3 is a restriction map of the vector pME18S.

Figures 4A and 4B illustrate the binding of 125 I-human 1L-4 to TF1 cells (A) and to COS 7 cells (B), where both types of cells are transiently transfected with pME18S-hIL-4R.

Figure 5 (in two parts A and B on two sheets) is a comparison of the amino acid sequences of human IL-4R (upper line of each pair) and mouse IL-4R (lower line of each pair), and shows the leader sequence of human IL-4R singly underlined and the transmembrane domain of human IL-4R doubly underlined.

DETAILED DESCRIPTION OF THE INVENTION

I. Obtaining and Expressing IL-4R cDNAs

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The term "effectively homologous" as used herein means that the nucleotide sequence is capable of being detected by a hybridization probe derived from a cDNA clone of the invention. The exact numerical measure of homology necessary to detect nucleic acids coding for an IL-4R depends on several factors including (1) the homology of the probe to sequences that do not encode it.-4R but are associated with the target nucleic acids, (2) the stringency of the hybridization conditions, (3) whether single-stranded or double-stranded probes are employed, (4) whether RNA or DNA probes are employed, (5) the measures taken to reduce nonspecific binding of the probe. (6) the nature of the method used to label the probe. (7) the fraction of guanidine and cytosine bases in the probe, (8) the distribution of mismatches between probe and target, (9) the size of the probe, and the like. Preferably, an effectively homologous nucleic acid sequence is at least seventy percent (70%) homologous to the cDNA of the invention. More preferably, an effectively homologous nucleic acid is at least ninety percent (90%) homologous to the cDNA of the invention. Most particularly, an effectively homologous nucleic acid sequence is one whose cDNA can be isolated by a probe based on the nucleic acid sequence of Formula I using the hybridization protocol described in the Examples with no more than a few false positive signals, e.g. fewer than a hundred. There is an extensive literature that provides guidance in selecting conditions for such hybridizations: e.g. Hames et al., Nucleic Acid Hybridization: A Practical Approach (IRL Press, Washington, D.C., 1985); Gray et al., Proc. Natl. Acad. Sci. USA, Vol. 80, pgs. 5842-5846 (1983); Kafatos et al., Nucleic Acids Research, Vol. 7. pgs. 1541-1552 (1979); and Williams, Genetic Engineering , Vol. 1, pgs. 1-59 (1981). By way of example, the nucleic acid of Formula I can be used as a probe in colony hybridization assays as described by Benton and Davis in Science, Vol. 196, pg. 180 (1977). Preferably, low stringency conditions are employed for the probe employed; the dissociation temperature depends on probe length, and increases as the probe length increases. For example, for a probe of about 20-40 bases a typical protocol including the steps of: prehybridizing, hybridizing, and washing; is as follows: (1) prehybridizing: incubate nitrocellulose filters containing the denatured target DNA for 3-4 hours at 55 °C in 5x Denhardt's solution, 5x SSPE (20x SSPE consists of 174 g NaCl, 27.6 g NaH₂PO₄.H₂O₄ and 7.4 g EDTA in 800 ml water adjusted to pH 7.4 with 10 N NaOH), 0.1% SDS, and 100 u.g/ml denatured salmon sperm DNA, (2) hybridizing: incubate filters in prehybridization solution plus probe at 55°C for 2 hours, (3) washing: three 15-minute washes in 300-500 ml volumes of 8x SSC and 0.1% SDS at room temperature, followed by a final 1-1.5-minute wash in 300-500 ml of 1x SSC and 0.1% SDS at 55°C. Other equivalent procedures are well known in the art.

Homology as the term is used herein is a measure of similarity between two nucleotide (or amino acid) sequences. Homology is expressed as the fraction or percentage of matching bases (or amino acids) after two sequences (possibly of unequal length) have been aligned. The term alignment is used in the sense defined by Sanfroff and Kruskal in Chapter 1 of "Time Warps, String Edits, and Macromolecules: The Theory and Practice of Sequence Comparison" (Addison-Wesley, Reading, MA, 1983). Roughly, two sequences are aligned by maximizing the number of matching bases (or amino acids) between the two sequences with the insertion of a minimal number of "blank" or "null" bases into either sequence to bring about the maximum overlap. Given two sequences, algorithms are available for computing their homology, e.g. Needleham and Wunsch, J. Mol. Biol., Vol. 48, pgs. 443-453 (1970), and Sankoff and Kruskal (cited above) pgs. 23-29. Also, commercial services and software packages are available for performing such comparisons, e.g. Intelligenetics, Inc. (Mountain View, CA), and University of Wisconsin Genetics Computer Group (Madison, Wisconsin).

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Probes based on the nucleic acid sequence of Formula I can be synthesized on commercially available DNA synthesizers, e.g. Apptied Biosystems model 381A, using standard techniques, e.g. Gait, Oligonucleotide Synthesis: A Practical Approach, (IRL Press, Washington D.C., 1984). Preferably the probe is at least 18-30 bases long, more preferably at least 100-200 bases long. Probes of the invention can be labeled in a variety of ways standard in the art; e.g. with radioactive labels (see e.g. Berent et al., Biotechniques, pgs. 208-220 (May/June 1985), Meinkoth et al., Anal. Biochem. Vol. 138, pgs. 287-284 (1984), and Szostak et al., Meth. Enzymol. Vol. 68, pgs. 419-429 (1979)), or with non-radioactive labels (see e.g. Chu et al., DNA, Vol. 4, pgs. 327-331 (1985), Jablonski et al., Nucleic Acids Research, Vol. 14, pgs. 6115-6128 (1986)).

Hybridization probes can also be used to screen candidate sources of IL-4R mRNA prior to library construction: e.g. by RNA blotting, Maniatis et al., Molecular Cloning: A Laboratory Manual, pgs. 202-203 (Cold Spring Harbor Laboratory, N.Y., 1982); or Harnes and Higgins, eds., pgs. 139-143 in Nucleic Acids Hybridization (IRL Press, Washington, D.C., 1985). Sources of mRNA encoding the desired polypeptides include cell populations or cell lines that express, or can be induced to express, large numbers of IL-4 receptors on their surfaces, e.g. in excess of 3000-5000. Such cells include T cells, T cell lines, mast call lines, and the like. In general, suitable T cells can be obtained from a variety of sources, such as mammalian (e.g. human) spleen, tonsils and peripheral blood. Preferably, mRNA is obtained from T cells extracted from the peripheral blood and treated with IL-4 to induce an increased expression of IL-4R. Such induction can be accomplished by first exposing the human T cells to 1 µg ml of phytohemaglutinin (PHA) for about 3 days, followed by exposure to about 1000 picomolar IL-4. After about 24 hours, the cells express a 2-3 fold increase in IL-4Rs.

If the mRNAs encoding a desired IL-4R make up a very small fraction of the total mRNA, it may be necessary to enrich the fractional concentration in order to make practical the screening procedure for detecting cDNA clones of interest. Such procedures are standard in the art and are disclosed in several papers of references, such as: Maniatis et al., pgs. 225-228, cited above; Suggs et al., Proc. Natl. Acad. Sci. USA, Vol. 78, pgs. 6813-6817 (1981); Pames et al., Proc. Natl. Acad. Sci. USA, Vol. 78, pgs. 2253-2257 (1981); Davis et al., Proc. Natl. Acad. Sci. USA, Vol. 81, pgs. 2194-2198 (1984); or the like.

Once a cDNA of the invention has been cloned, a wide range of expression systems (i.e. hostexpression vector combinations) can be used to produce the proteins of the invention. Possible types of 30 host cells include, but are not limited to, bacterial, yeast, insect, mammalian, and the like. Many reviews are available which provide guidance for making choices and/or modifications of specific expression systems: e.g. de Boer and Shepard, "Strategies for Optimizing Foreign Gene Expression in Escherichia coli." pgs. 205-247 in Kroon, ed. Genes: Structure and Expression (John Wiley & Sons, New York, 1983), review several E. coli expression systems; Kucherlapati et al., Critical Reviews in Biochemistry , Vol. 16, Issue 4, pgs. 348-379 (1984); and Banerji et al., Genetic Engineering , Vol. 5, pgs. 19-31 (1983) review methods for transfecting and transforming mammalian cells; Reznikoff and Gold, eds., Maximizing Gene Expression (Butterworths, Boston, 1986) review selected topics in gene expression in E. coli , yeast, and mammalian cells; and Thilly, Mammalian Cell Technology (Butterworths, Boston, 1986) reviews mammalian expression systems (to mention a few). Likewise, many reviews are available which describe techniques and conditions 40 for linking and/or manipulating specific cDNAS and expression control sequences to create and/or modify expression vectors suitable for use with the present invention and to create and modify cDNAS to produce soluble forms of the receptor: e.g. Maniatis et al., Molecular Cloning: A Laboratory Manual (Cold Spring Harbor Laboratory, N.Y., 1982); Glover, DNA Cloning: A Practical Approach, Vol. I and II (IRL Press, Oxic d, 1985); and Perbal, A Practical Guide to Molecular Cloning (John Wiley & 50ns. N.Y., 1984), to name only a

An E. coli expression system is disclosed by Riggs in U.S. Patent 4,431,739, which is incorporated by reference. A particularly useful prolaryotic promoter for high expression in E. coli is the tac promoter, disclosed by de Boer in U.S. Patent 4,551,433, which is incorporated herein by reference. Secretion expression vectors are also available for E. coli hosts. Particularly useful are the plN-III-ompA vectors, disclosed by Ghraveb et al. in EMBO J., Vol. 3, pgs. 2437-2442 (1994), in which the cDNA to be transcribed is fused to the portion of the E. coli OmpA gene encoding the signal peptide of the ompA protein which, in turn, causes the mature protein to be secreted into the periplasmic space of the bacteria. U.S. Patents 4,338,336 and 4,338,397 also disclose secretion expression vectors for prokaryotes and accordingly are incorporated by reference. Numerous strains of bacteria are suitable hosts for prokaryotic expression vectors including strains of E. coli, such as W3110 (ATCC No. 27325), JA221, C800, ED767, DH1, LE392, HB101, X1778 (ATCC No. 31244), X2282, RR1 (ATCC No. 31343) MRCI; strains of Bacillus subtilis; and other enterobac teriaceae such as Salmonella typhimurium or Serratia marcescens, and various species of Pseudomonas. General methods for deriving bacterial strains, such as E. coli K12

X1776, useful in the expression of eukaryotic proteins are disclosed by Curtis III in U.S. Patent 4,190,495, which is accordingly incorporated by reference.

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In addition to prokaryotic and eukaryotic microorganisms, expression systems comprising cells derived from multicellular organisms may also be used to produce proteins of the invention. Of particular interest are mammalian expression systems because their posttranslational processing machinery is more likely to produce biologically active mammalian proteins. Several DNA tumor viruses have been used as vectors for mammalian hosts. Particularly important are the numerous vectors which comprise SV40 replication, transcription, and/or translation control sequences coupled to bacterial replication control sequences: e.g. the pcD vectors developed by Okayama and Berg, disclosed in Mol. Cell. Biol., Vol. 2, pgs. 161-179(1982) and Mol. Cell. Biol., Vol. 3, pgs. 280-289 (1983), both of which are incorporated herein by reference; the SV40 vectors disclosed by Hamer in Genetic Engineering. Vol. 2, pgs. 83-100 (1980), and U.S. Patent 4,599,308, both of which are incorporated herein by reference; and the vectors additionally containing adenovirus regulatory elements, disclosed by Kaufman and Sharp, in Mol. Cell. Biol., Vol. 2, pgs. 1304-1319 (1982), and Clark et al., in U.S. patent 4,675,285, both of which are incorporated herein by reference.

COS7 monkey cells, described by Gluzman, Cell., Vol. 23, pgs. 175-182 (1981) and available from the ATCC (accession no. CRL 1651), are usually the preferred hosts for the above vectors.

Preferably, soluble forms of the IL-4Rs are produced by introducing a stop codon prior to (i.e. in the 5-or "upstream" direction of) the coding region for the transmembrane and intracellular politions of the IL-4Rs cDNA. This is conveniently done by site-specific mutagenesis, e.g. as taught by Zoller and Smith, Meth. Enzymol., Vol. 100, pgs. 468-500 (1983), and the like. Extracellular, transmembrane, and intracellular domains of receptor cDNAs are readily identified by the hydrophobicity of the encoded amino acids in the largest open reading frame. For example, the transmembrane domain is a segment containing from about 20-25 residues having high average hydrophobicity, e.g. Wickner, Science, Vol. 210, pgs. 881-888 (1980), and Greene et al., Ann. Rev. Immunol., Vol. 4, pgs. 68-95 (1986). The amino acids in the N-terminal direction from the transmembrane domain constitute the extracellular domain, and the amino acids in the C-terminal direction from the transmembrane domain constitute the intracellular domain.

II. Soluble Interleukin-4 Receptor Binding Assays

The affinity, or strength of binding, of soluble IL-4Rs to IL-4 is measured in relation to the binding of IL-4 to a standardized number of membrane-bound IL-4Rs. That is, the affinity of a soluble IL-4R for IL-4 is determined in a competition assay with membrane-bound IL-4R. The membrane-bound IL-4Rs can be the natural IL-4Rs of a cell line responsive to IL-4, e.g. a helper T cell line. Affinity can be measured in terms of inhibition of biological activity or the inhibition of binding of 125 Habeled IL-4 to the membrane-bound IL-4Rs.

Typically the biological activity measured is cell growth, as determined by tritiated thymidine incorporation, or like assays of cell growth, e.g. Mosmann, J. Immunol. Meth., Vol. 65, pgs. 55-63 (1983). Thus, inhibition of IL-4-induced cell growth is measured by exposing each of a series of cell cultures to a constant initial concentration of IL-4 and a different initial concentration of soluble IL-4R, and then examining the relationship of soluble IL-4R concentration to cell growth., e.g. as illustrated by Figure 2. Preferably, a cellline responsive to IL-4 is used, but T cells from peripheral blood lymphocytes can also be used. Preferably, the concentration of IL-4 is at or near the level where the biological response becomes saturated, i.e. the level where additional IL-4 has no measurable effect on the cells. Accordingly, to apply the inhibition at say for particular cell types may require some routine experimentation to establish the saturation concentration. By way of example, inhibition of human T cell growth factor activity can be assayed by the following steps: (1) aliquots of washed human peripheral blood lymphocytes (about 2 x 10^5 in 50 μ I) previously stimulated with phytohemaggiutinin (PHA) for 7 days and subsequently cultured for 7 days with IL-2 are added to microtiter plates each containing a known concentration of IL-4, (2) dilutions (e.g. in 50 µl) of the stL-4R are added to each well, (3) the cells are incubated for 72 hours at 37°C. (4) tritiated thyrnidine (about 20 ml at 20 µCi/ml) is added to each well, and (5) cells are harvested onto filter paper, washed, and counted in a scintillation counter.

For the alternative assay, IL-4 is iodinated by standard commercially available techniques, e.g. reaction with aqueous Na¹²⁵1 and with 1,3,4,6-tetrachloro-3a,6a-diphenylglycoluril (described by Fraker et a]. Biochem Biophys. Res. Commun., Vol. 80, pgs. 849-857 (1978) and available from Pierce Chemical Co. as IODO-GEN). Generally, the binding assay is as described by Lowenthal et al., J. Immunol., Vol. 140, pgs. 458-464 (1988), which is incorporated by reference. Human cell lines having IL-4R suitable for use with the binding assays include Jiyoye cells and Jurkat cells, the former being available from the American Type Culture Collection under accession number CCL 87.

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III. Purification and Pharmaceutical Compositions

When polypeptides of the present invention are expressed in soluble form, for example as a secreted product of transformed yeast or mammalian cells, they can be purified according to standard procedures of the art, including steps of ammonium sulfate precipitation, ion exchange chromatography, gel filtration, electrophoresis, affinity chromatography, and/or the like: e.g. "Enzyme Purification and Related Techniques," Methods in Enzymology . 22:233-577 (1977), and Scopes. R., Protein Purification: Principles and Practice, Springer-Verlag, New York [1982]) provide guidance in such purifications. Likewise, when polypeptides of the invention are expressed in insoluble form, for example as aggregates, inclusion bodies or the like, they can be purified by standard procedures in the art, including separating the inclusion bodies from disrupted host cells by centrifugation, solubilizing the inclusion bodies with chaotropic agents and reducing agents, diluting the solubilized mixture, and lowering the concentration of chaotropic agent and reducing agent so that the polypeptide takes on a biologically active conformation. The latter procedures are disclosed in the following references, which are incorporated by reference: Winkler et al., Biochemistry . 25: 4041-4045 (1988); Winkler et al., Biotechnology . 3: 992-998 (1985); Koths et al., U.S. patent 4.569,790; and European patent applications 86308917.5 and 86308353.3.

For preparing pharmaceutical compositions containing the polypeptides described by this invention, these polypeptides are compounded by admixture with preferably inert, pharmaceutically acceptable carriers. Suitable carriers and processes for their preparation are well known in the art. e.g., Remington's Pharmaceutical Sciences and U.S. Pharmacopeia: National Formulary (Mack Publishing Company, Easton, PA,1984). The preferred course of administration is parenteral and can include use of an implantable drug delivery system, e.g. Urguhart et al., Ann. Rev. Pharmacol. Toxicol., Vol. 24, pgs. 199-236 (1984).

As used herein "effective amount" means an amount sufficient to ameliorate a symptom of an autoimmune condition. The effective amount for a particular patient may vary depending on such factors as the state of he autoimmune condition being treated, the overall health of the patient, method of administration, the severity of side-effects, and the like. Generally, IL-4R is administered as a pharmaceutical composition comprising an effective amount of IL-4P and a pharmaceutical carrier. A pharmaceutical carrier can be any compatible, non-toxic substance suitable for delivering the compositions of the invention to a patient. When administered parenterally, the IL-4R is formulated in a unit dosage injectable form (e.g., 30 solution, suspension, or emulsion) in association with a pharmaceutical carrier. Such carriers are inherently nontherapeutic and nontoxic. Examples of such carriers are normal saline, Ringer's solution, dextrose solution, and Hank's solution. Nonaqueous carriers such as fixed oils and ethyl cleate may also be used. A preferred carrier is 5% dextrose saline. The carrier may contain minor amounts of additives such as substances that enhance isotonicity and chemical stability, e.g., buffers and preservatives. The IL-4R is 35 preferably formulated in purified form substantially free of aggregates and other proteins at a concentration in the range of about 5 to 20 u.g.ml. Preferably, IL-4R is administered by continuous infusion so that an amount in the range of about 50-800 µg is delivered per day (i.e. about 1-16 µg/kg/day). The daily infusion rate may be varied based on monitoring of side effects and IL-4 and/or IgE levels.

Preferably, IL-4R is purified from culture supernatants of mammalian cells transiently transfected or 40 stably transformed by an expression vector carrying an IL-4R gene. Most preferably, IL-4R is purified from culture supernatants of COS 7 cells transiently transfected by the pcD expression vector. Transfection of COS 7 cells with pcD proceeds as follows: One day prior to transfection, approximately 106 COS 7 monkey cells are seeded onto individual 100 mm plates in Dulbecco's modified Eagle medium (DME) containing 10% fetal call serum and 2 mM glutamine. To perform the transfection, the medium is aspirated from each 45 plate, washed with DME containing 50 mM Tris.HCl pH 7.4, and replaced with 4 ml of DME containing 50 mM Tris.HCl pH 7.4, 400 µg/ml DEAE-Dextran and 5 µg of plasmid DNA. The plates are incubated for four hours at 37°C, and then the DNA-containing medium is removed, and the plates are washed once with 5 ml of PBS, and then DME containing 10% fetal calf serum, 2 mM glutamine, penicillin and streptomycin is added. The cells are then incubated for 72 hrs at 37°C, after which the growth medium is collected for 50 purification of IL-4R. Alternatively, transfection can be accomplished by electroporation as described in the Examples. Plasmid DNA for the transfections is obtained by growing pcD(SRa) containing the IL-4R cDNA insert in E. coli MC1081, described by Casadaban and Cohen, J. Mol. Biol., Vol. 138, pgs. 179-207 (1980), or like organism. The plasmid DNA is isolated from the cultures by standard techniques, e.g. Maniatis et al., Molecular Cloning: A Laboratory Manual (Cold Spring Harbor Laboratory, New York, 1982).

EXAMPLES

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Example I. Construction of cDNA library from MC-9 cells and isolation of clone 19

A cDNA library was constructed in the pCEV4 vector from mRNA extracted from MC.9 cells, disclosed in U.S. patent 4,559,310, pCEV4 pcD and its derivative, pcD(SRa), are described by Okayama and Berg, Mol. Cell. Biol., Vol. 2: 161-170 (1982) and Vol. 3: 280-289 (1983), and by Takebe et al., Mol. Cell. Biol., Vol. 8, pgs. 466-472 (1988), respectively. The polyoma virus origin (the Bgll-Bcll fragment) was inserted into the Nde I site of pcD(SRa) for replication in muuse cells, and a polylinker sequence containing two Bst XI sites was inserted in the Pst I site of pcD(SRa). Size-selected double stranded cDNAS (>1.5 kilobases) having Bat XI adaptors were inserted into Bat XI-digested pCEV4, amplified in E. coli , and used to transfect 10 COS 7 monkey cells by protoplast fusion (e.g. Sandri-Goldin et al., Mol. Cell. Biol., Vol. 1, pgs. 743-752. (1981)). Populations of COS cells enriched for those expressing IL-4R on their surfaces were obtained by a modified panning technique based on that described by Seed (e.g. Proc. Natl. Acad. Sci. USA . Vol. 84. pgs. 3365-3369 (1987), or Proc. Natl. Acad. Sci. USA , Vol. 84. pgs 8573-8577 (1987)). Briefly, the modification consisted of the following steps: (1) biotin was attached to lysine residues of mouse IL-4. (2) is prior to panning, the biotinylated IL-4 was incubated with the transfected COS cells so that the biotinylated IL-4 could bind to IL-4Rs expressed on the COS cells, and (3) the bound biotinylated IL-4 was covalently cross linked to the receptors by bis(2-(succinimidooxycarbonyloxy)ethyl)sulfone (Pierce Chemical Company. Rockford, IL). After these initial three steps the standard panning protocol was employed using microtitar plates coated with an anti-biotin antibody to selectively bind COS cells expressing IL-4Rs. Panning resulted in the isolation of clone 19 carrying the cDNA whose sequence is listed in Formula I.

Example II. Inhibition of IL-4 binding to T cells by soluble IL-4R

A cDNN encoding a soluble It.-4R was constructed from the It.-4R cDNA isolated in Example 1 by inserting linker CTAGACTAGTCTAG into a unique Bst Ell restriction endonuclease site of clone 19 that was adjacent to and just 5 - of the transmembrane encoding domain. The resulting soluble It.-4R consisted of the extracellular domain of the receptor encoded by clone 19. The modified cDNA encoding the soluble It.-4R was transfected into COS 7 cells and the culture supernatants were assayed for inhibition of ¹²⁵Habeled It.-4 binding to the It.-4Rs of the mouse T cell line, HT-2; and for the inhibition of It.-4-induced growth of HT-2. The results are illustrated in Figures 1 and 2, respectively. In the growth inhibition assay the concentration of It.-4 was 100 picomolar.

Example III. Use of clone 19 to screen for a human iL-4R cDNA

Messenger RNA was prepared from the human Burkitt lymphoma cell line, BL41, and the human multi-factor dependent myeloid cell line, TF1, by the guanidium thiocyanate method. RNA was converted to cDNA, and Bst XI adaptors were attached to the cDNA cDNAS larger than 2.0 kilobases were isolated by electrophoresis on a 0.8% agarose gel. The cDNA library of BL41 was constructed by inserting the size-selected cDNA into the Bst XI sites of the pCEV4 vector, a derivative of the pcD(SRa) expression vector described by Itoh et al., Science, Vol. 247, pg. 324 (1990), 1.2 x 10⁶ independent clones were obtained. A kgt11 phage cDNA library of TF1 was constructed by using the oligo(dT) primer containing the Not restriction site and the Eco RI adaptor according to the protocol provided by Promega Corp. (Madison, WI).

The BL41 cDNA library was screened by colony hybridization using the mouse IL-4R cDNA as a probe. Colonies were transferred to nylon membrane filters (Schleicher & Schuell), and hybridization was performed at 42°C in 6 x SSPE (1 x SSPE contains 17.4 g l NaCl, 2.78 g/l NaH₂PO₄.H₂O, 7.4 g/l EDTA, pH 7.4) containing 20% formamide, 5X Denhardt's solution, and 100 mg/ml E. coli tRNA. Filters were washed with 4 x SSPE solution at 50°C and exposed to X-ray film. Partial nucleotide sequence analysis showed that the longest cDNA clone from the BL41 library (4A2) had approximately 70 percent homology with the mouse IL-4R, but lacked about 700 bases from the 5 end. 4A2 was then used as a probe to screen the TFI cDNA library under high stringency conditions (hybridization in 50% formamide at 42°C, and wash in 0.2 x SSPE at 60°C, otherwise the same as above). A clone (TFI-2.2) was obtained containing a 3.6 kilobase cDNA insert.

The nucleotide sequence of the 3.6 kilobase insert of the human IL-4R cDNA revealed a single long oper, reading frame encoding 825 amino acid residues including a putative signal sequence of 25 amino acid residues (indicated by underlining in Figure 5), an extracellular domain of 207 amino acid residues, a transmembrane domain (indicated by double underlining in Figure 5), and a cytoplasmic domain of 569

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amino acid residues.

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This soluble form of the human interleukin-4 receptor consists of an extracellular domain of the human interleukin-4 receptor. The extracellular domain typically consists of 200-220 N-terminal amino acids of the human interleukin-4 receptor. The 207-unit amino acid sequence prepared in the foregoing Example has the following amino acid sequence:

MKVLQEPTCV	SDYMSISTCE	WKMNGPTNCS	TELRLLYQLV
FLLSEAHTCV	PENNGGAGCV	CHLLMDDVVS	ADNYTLDLWA
GQQLLWKGSF	KPSEHVKPRA	PGNLTVHTNV	SDTLLLTWSN
PYPPDNYLYN	HLTYAVNIWS	ENDPADFRIY	NVTYLEPSLR
IAASTLKSGI	SYRARVRAWA	QCYNTTWSEW	SPSTKWHNSY
REPFEOH:			

where the amino acids are designated by their single-letter abbreviations, namely

A	Ala	Alanine	М	Met	Methionine
С	Cys	Cysteine	N	Asn	Asparagine
D	Aso	Aspartic acid	P	Pro	Proline
Ε	Glu	Glutamic acid	Q	Gin	Glutamine
F	Phe	Phenylalanine	R	Arg	Arginine
G	Gly	Glycine	S	Ser	Serine
н	His	Histidine	T	Thr	Threonine
1	lle	Iscleucine	V	Vai	Valine
ĸ	Lys	Lysine	W	Trp	Tryptophane
L	Leu	Leucine	Y	Tyr	Tyrosine.

Example IV. Binding properties of the human IL-4R

The full length human IL-4R cDNA was inserted into the mammalian expression vector, pME18S (Figure 3), and was transfected into COS 7 cells. pME18S-hIL-4R is deposited with the American Type Culture Collection (Rockville, MD) under accession number 68283. Binding assays were carried out as follows on the transfected COS 7 cells and the TF1 cells. COS 7 cells were detached from the plates by a short incubation in PBS containing 0.5 mM EDTA. Exponentially growing TF1 cells were harvested by centrifugation. Cells were washed and resuspended in RPMI 1640 containing 20 mM HEPES, and 0.5% BSA. Cells (7 x 10° COS 7 cells or 1.2 x 10° TF1 cells) were incubated with 5-250 pM of '25 I-IL-4 in the presence (solid squares or circles) and absence (empty squares or circles) of 10 nM nonradioactive IL-4 at + ° C for 3 hours. To prevent cell loss during washing, 10° mouse MC-9 cells, which do not bind human IL-4, were added to each sample at the end of the incubation. Cell-bound radioactivity was separated by centrifugation as described by Galizzi et al., J. Biol. Chem., Vol. 264, pg. 6864 (1989), which reference is incorporated by reference. The affinity of IL-4 binding to its receptor and the number of IL-4 receptors per cell were estimated by Scatchard plot analysis of equilibrium binding using the LIGAND program disclosed by Munson, Meth. Enzymol., Vol. 92, pgs. 543-578 (1983).

Equilibrium binding of ¹²⁵I-human IL-4 to the translected COS 7 cells and to the TF1 cells showed a single class of high affinity binding sites in both cells (Figure 4). Scatchard analysis indicated that COS 7 cells expressed 140,000 sites/cell with a binding constant, K_d, of 80-100 pM, while TF1 cells expressed about 1000 receptors/cell with a K_d of 40-60 pM (Figure 4).

The descriptions of the foregoing embodiments of the invention have been presented for purpose of illustration and description. They are not intended to be exhaustive or to limit the invention to the precise forms disclosed, and obviously many modifications and variations are possible in light of the above teaching. The embodiments were chosen and described in order to best explain the principles of the

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invention to thereby enable others skilled in the art to best utilize the invention in various embodiments and with various modifications as are suited to the particular use contemplated. It is intended that the scope of the invention be defined by the claims appended hereto.

Applicants deposited E. coli MC1081 carrying pME18S-hIL-4R with the American Type Culture Collection, Rockville, MD, USA (ATCC), under accession number 68283, on March 20th 1990. This deposit was made under conditions as provided under ATCC's agreement for Culture Deposit for Patent Purposes, which assures that the deposit will be made available to the US Commissioner of Patents and Trademarks pursuant to 35 USC 122 and 37 CFR 1.14, and will be made available to the public upon issue of a U.S. patent, which requires that the deposit be maintained. Availability of the deposited strain is not to be construed as a license to practise the invention in contravention of the rights granted under the authority of any government in accordance with its patent laws.

The Deposits have been modified to comply with the requirements of the Budapest Treaty on the Deposit of Microorganisms.

Claims

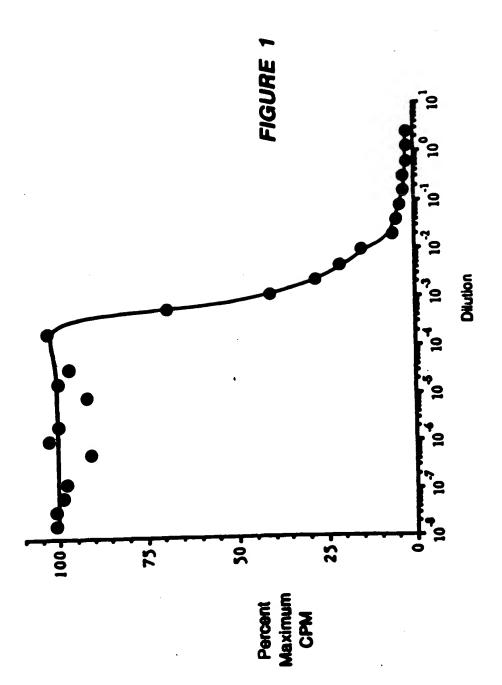
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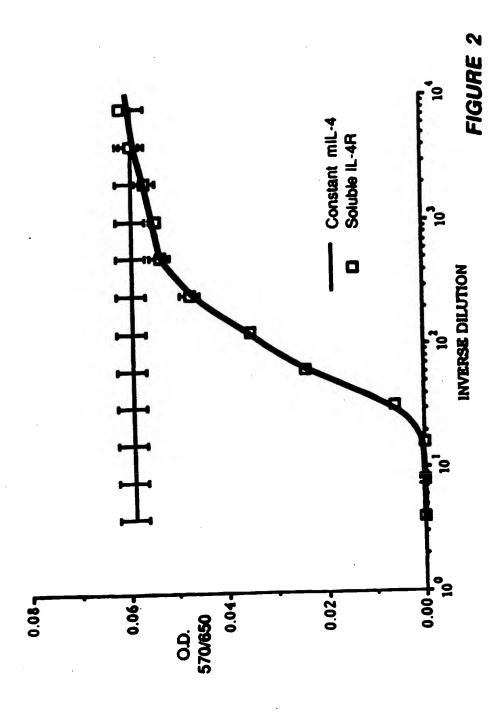
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- 1. A nucleic acid encoding a mammalian interleukin-4 receptor.
- 2. The nucleic acid of claim 1 wherein said mammalian interleukin-4 receptor is a human interleukin-4 receptor.
 - 3. The nucleic acid of claim 2 encoding a soluble form of said human interleukin-4 receptor.
 - 4. The nucleic acid of claim 3 wherein said soluble form of said human interleukin-4 receptor consists of an extracellular domain of said human interleukin-4 receptor.
- The nucleic acid of claim 4 wherein said extracellular domain consists of the 200-220 N-terminal aminoacids of a id human interleutin-4 receptor.
 - 6. The nucleic acid of claim 2 which is effectively homologous to the nucleotide sequence of Formula 1 herein.
 - 7. A soluble human interleukin-4 receptor consisting of the extracellular domain of the human interleukin-4 receptor.
- 30 8. The soluble human interleukin-4 receptor of claim 7 having the following amino acid sequence:

MKVLQEPTCV	SDYMSISTCE	WKMNGPINCS	TELRLLYQLV
FLLSEAHTCV	PENNGGAGCV	CHLLMDDVVS	AWIGITYNDA
GQQLLWKGSF	KPSEHVKPRA	PGNLTVHTNV	SDTLLLTWSN
PYPPDNYLYN	HLTYAVNIWS	ENDPADFRIY	NVTYLEPSLR
IAASTLKSGI	SYRARVRAWA	QCYNTTWSEW	SPSTKWHNSY
REPFEQH.			

9. Plasmid pME188-htt-4R deposited at the ATTC under accession no 68263.





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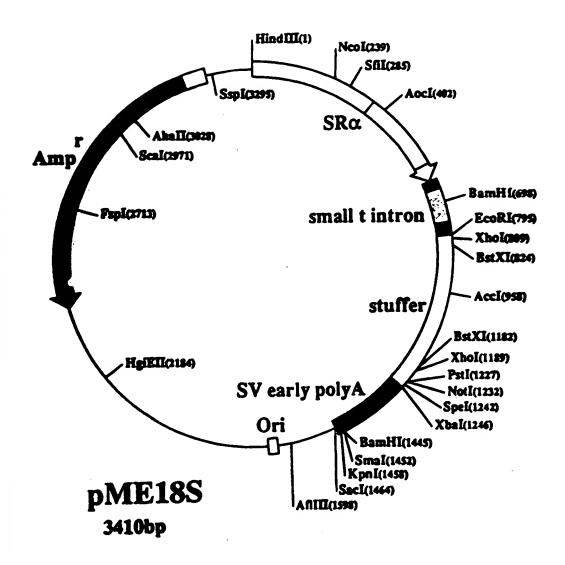


FIGURE 3

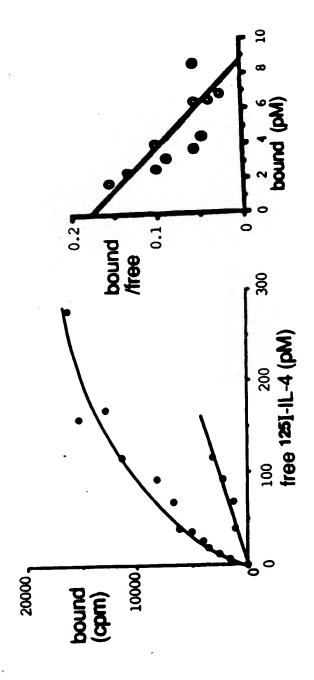


FIGURE 4 part A

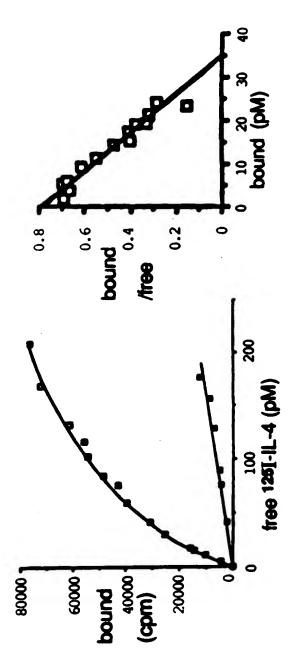


FIGURE 4 part B

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1	MGwLCsglLfpVsCLvLLqVasSGnmKVLqEPTCMGrLCtkfLtsVgCLiLLlVtgSGsiKVLgEPTC	• • •
35 35	vSDYmsiSTCEWkmngptnCS-telrllyqlvFlfSDYirtSTCEWfldsandCSsqlclhyrlmfFe	
68 69	1SEahTCvPeNnggagCVCHllMddvVsaDnYtl fSEnlTCiPrNsastvCVCHmeMnrpVqsDrYqm	1
102 103	dLWAgqqlLWkGSFkPSehVKPrAPgNLTvHTNVeLWAehrqLWqGSFsPSgnVKPlAPdNLTlHTNV	7 7
136 137	SDtlLLTWsNpYPpdNyLYnhLtyaVNIwsEndP SDewLLTWnNlYPsnNlLYkdLismVNIsrEdnP)
170 171	AdFriYNVTYlEPsLriaastLkSGisYrARVRa AeFivYNVTYkEPrLsfpiniLmSGvyYtARVR	<u>,</u>
204 205	waQcyntTWSEWSPStkWhNsyrePfeQhLlLGVrsQiltgTWSEWSPSitWyNhfqlPliQrLpLGV	<u>/</u>
238 239	SVSCIVILAVCLICYVSITKIKKEWWDQIPnPAF tiscicipiicticyisitkikkiwwdQIPtPAF	ર ર
272 273	SrlvailiQDAQgsqWeKrsRgQEpaKcPHWKnCSplvailiQDAQvplWdKqtRsQEstKyPHWKtC	777
306 307	LtKLLPCfLeHnmKrdeDphKAAkempfQgsGKs LdKLLPC1LkHrvKkktDfpKAAptks1QspGKa	5 3
340 341	aWCPvEiSkTVLWPESiSVVRCvELFEAPV-egWCPmEvSrTVLWPEnvSvSVVRCmELFEAPVqr	e
371 375	cEEEeveeekgsfcaspeSsrddFQEgregIvivEEEEdeivkedlsmspenSggcgFQEsqadImi	A A
405 409	RLTESLF1DLLgeENGGfcQqdmgESC111PSG RLTEnLFSDLLeaENGG1gQsalaESCsp1PSG	s s

FIGURE 5 part A

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439 443	tsAhmpWdefPsagpkEAppwgkEQPlhlepsppgqAsvsWaclPmgpseEAtcqvtEQP-shpgpls	
473 476	aSPtQSpdnLtCTetPLViAgNPAYRSFSnslSq gSPaQSaptLaCTqvPLVlAdNPAYRSFSdccSp	
507 510	sPcPrELgPdpllArHLEEvEPemPcvpqlseptaPnPgELaPeqqqAdHLEEeEPPspadphssg	
541 542	t vpQPepEtWE QILrrnVLQHGAAAapvsAPtsG p pmQP-vEsWE QILhmsVLQHGAAAgstpAPagG	
575 575	YQEFVhAVeQGgtQasaVvGlgPpGeaGYKAFSS YQEFVqAVkQGaaQdpgVpGvrPsGdpGYKAFSS	
609 609	LLaSsavspekcgfGassGeeGYKPFQdlipgcp LLsSngirgdtaaaGtddGhgGYKPFQnpv	
643 639	gdpaPvpVPLFTFGLDrEpprSPqsShlPsSsPE p	
677 673	hLGLEpGeKveDmpKpPlPqeQatdPlvDsLGsG cLGLElGlKggDwvKaPpPadQvpkPfgDdLGfG	
711 707	IVYSaLTCHLCGHLKQcHgQEdGGQtPvmASPcC IVYSsLTCHLCGHLKQhHsQEeGGQsPivASPgC	
745 741	GCCcgDRSsppttplrApdpsPgGvPLEAslcpa GCCydDRSpslgslsgAlescPeGiPLEAnl	
779 772	sl APsgisekskss SsfhpaPGnaqssSQTpkiv m sAP ktpsnlSgegkgPGhspvpSQT	
813 798	nfVsVGptymrVS 825 teVpVGalgiaVS 810	

FIGURE 5 part B



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Application Number

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Citation of document with indication, where appropriate. Relevant			CLASSIFICATION OF THE	
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